

Revised

IFWO

RAW SEQUENCE LISTING

DATE: 10/15/2004

PATENT APPLICATION: US/10/734,661A

TIME: 11:07:01

Input Set : N:\AMC\US10734661A.raw

Output Set: N:\CRF4\10152004\J734661A.raw

1 <110> APPLICANT: ProChon Biotech, Ltd.

2 MorphoSys AG

3 Yayon, Avner

4 Thomassen-Wolf, Elisabeth

5 Rom, Eran

6 Borges, Eric

7 <120> TITLE OF INVENTION: ANTIBODIES THAT BLOCK RECEPTOR PROTEIN TYROSINE KINASE

ACTIVATION

8 <130> FILE REFERENCE: 81408-4400

C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/734,661A

10 <141> CURRENT FILING DATE: 2003-12-15

11 <150> PRIOR APPLICATION NUMBER: US 60/299,187

12 <151> PRIOR FILING DATE: 2001-06-20

13 <150> PRIOR APPLICATION NUMBER: PCT/IL02/00494

14 <151> PRIOR FILING DATE: 2002-06-20

15 <160> NUMBER OF SEQ ID NOS: 106

16 <170> SOFTWARE: PatentIn version 3.2

18 <210> SEQ ID NO: 1

19 <211> LENGTH: 806

20 <212> TYPE: PRT

21 <213> ORGANISM: Homo sapiens

22 <300> PUBLICATION INFORMATION:

23 <308> DATABASE ACCESSION NO: np_000133

24 <309> DATABASE ENTRY DATE: 2001-02-21

25 <313> RELEVANT RESIDUES: (1)..(806)

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31 Gly Arg Ala Ala Glu Val Pro Gly Pro Glu Pro Gly Gln Gln Glu Gln

32 35 40 45

33 Leu Val Phe Gly Ser Gly Asp Ala Val Glu Leu Ser Cys Pro Pro Pro

34 50 55 60

35 Gly Gly Gly Pro Met Gly Pro Thr Val Trp Val Lys Asp Gly Thr Gly

36 65 70 75 80

37 Leu Val Pro Ser Glu Arg Val Leu Val Gly Pro Gln Arg Leu Gln Val

38 85 90 95

39 Leu Asn Ala Ser His Glu Asp Ser Gly Ala Tyr Ser Cys Arg Gln Arg

40 100 105 110

41 Leu Thr Gln Arg Val Leu Cys His Phe Ser Val Arg Val Thr Asp Ala

42 115 120 125

43 Pro Ser Ser Gly Asp Asp Glu Asp Gly Glu Asp Glu Ala Glu Asp Thr

44 130 135 140

*f.6*

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45 Gly Val Asp Thr Gly Ala Pro Tyr Trp Thr Arg Pro Glu Arg Met Asp
46 145 150 155 160
47 Lys Lys Leu Leu Ala Val Pro Ala Ala Asn Thr Val Arg Phe Arg Cys
48 165 170 175
49 Pro Ala Ala Gly Asn Pro Thr Pro Ser Ile Ser Trp Leu Lys Asn Gly
50 180 185 190
51 Arg Glu Phe Arg Gly Glu His Arg Ile Gly Gly Ile Lys Leu Arg His
52 195 200 205
53 Gln Gln Trp Ser Leu Val Met Glu Ser Val Val Pro Ser Asp Arg Gly
54 210 215 220
55 Asn Tyr Thr Cys Val Val Glu Asn Lys Phe Gly Ser Ile Arg Gln Thr
56 225 230 235 240
57 Tyr Thr Leu Asp Val Leu Glu Arg Ser Pro His Arg Pro Ile Leu Gln
58 245 250 255
59 Ala Gly Leu Pro Ala Asn Gln Thr Ala Val Leu Gly Ser Asp Val Glu
60 260 265 270
61 Phe His Cys Lys Val Tyr Ser Asp Ala Gln Pro His Ile Gln Trp Leu
62 275 280 285
63 Lys His Val Glu Val Asn Gly Ser Lys Val Gly Pro Asp Gly Thr Pro
64 290 295 300
65 Tyr Val Thr Val Leu Lys Thr Ala Gly Ala Asn Thr Thr Asp Lys Glu
66 305 310 315 320
67 Leu Glu Val Leu Ser Leu His Asn Val Thr Phe Glu Asp Ala Gly Glu
68 325 330 335
69 Tyr Thr Cys Leu Ala Gly Asn Ser Ile Gly Phe Ser His His Ser Ala
70 340 345 350
71 Trp Leu Val Val Leu Pro Ala Glu Glu Glu Leu Val Glu Ala Asp Glu
72 355 360 365
73 Ala Gly Ser Val Tyr Ala Gly Ile Leu Ser Tyr Gly Val Gly Phe Phe
74 370 375 380
75 Leu Phe Ile Leu Val Val Ala Ala Val Thr Leu Cys Arg Leu Arg Ser
76 385 390 395 400
77 Pro Pro Lys Lys Gly Leu Gly Ser Pro Thr Val His Lys Ile Ser Arg
78 405 410 415
79 Phe Pro Leu Lys Arg Gln Val Ser Leu Glu Ser Asn Ala Ser Met Ser
80 420 425 430
81 Ser Asn Thr Pro Leu Val Arg Ile Ala Arg Leu Ser Ser Gly Glu Gly
82 435 440 445
83 Pro Thr Leu Ala Asn Val Ser Glu Leu Glu Leu Pro Ala Asp Pro Lys
84 450 455 460
85 Trp Glu Leu Ser Arg Ala Arg Leu Thr Leu Gly Lys Pro Leu Gly Glu
86 465 470 475 480
87 Gly Cys Phe Gly Gln Val Val Met Ala Glu Ala Ile Gly Ile Asp Lys
88 485 490 495
89 Asp Arg Ala Ala Lys Pro Val Thr Val Ala Val Lys Met Leu Lys Asp
90 500 505 510
91 Asp Ala Thr Asp Lys Asp Leu Ser Asp Leu Val Ser Glu Met Glu Met
92 515 520 525
93 Met Lys Met Ile Gly Lys His Lys Asn Ile Ile Asn Leu Leu Gly Ala

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95 Cys Thr Gln Gly Gly Pro Leu Tyr Val Leu Val Glu Tyr Ala Ala Lys
96 545          550          555          560
97 Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Asp
98          565          570          575
99 Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu Glu Gln Leu Thr Phe Lys
100          580          585          590
101 Asp Leu Val Ser Cys Ala Tyr Gln Val Ala Arg Gly Met Glu Tyr Leu
102          595          600          605
103 Ala Ser Gln Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Val Leu
104 610          615          620
105 Val Thr Glu Asp Asn Val Met Lys Ile Ala Asp Phe Gly Leu Ala Arg
106 625          630          635          640
107 Asp Val His Asn Leu Asp Tyr Tyr Lys Lys Thr Thr Asn Gly Arg Leu
108          645          650          655
109 Pro Val Lys Trp Met Ala Pro Glu Ala Leu Phe Asp Arg Val Tyr Thr
110          660          665          670
111 His Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe
112          675          680          685
113 Thr Leu Gly Gly Ser Pro Tyr Pro Gly Ile Pro Val Glu Glu Leu Phe
114 690          695          700
115 Lys Leu Leu Lys Glu Gly His Arg Met Asp Lys Pro Ala Asn Cys Thr
116 705          710          715          720
117 His Asp Leu Tyr Met Ile Met Arg Glu Cys Trp His Ala Ala Pro Ser
118          725          730          735
119 Gln Arg Pro Thr Phe Lys Gln Leu Val Glu Asp Leu Asp Arg Val Leu
120          740          745          750
121 Thr Val Thr Ser Thr Asp Glu Tyr Leu Asp Leu Ser Ala Pro Phe Glu
122          755          760          765
123 Gln Tyr Ser Pro Gly Gly Gln Asp Thr Pro Ser Ser Ser Ser Gly
124 770          775          780
125 Asp Asp Ser Val Phe Ala His Asp Leu Leu Pro Pro Ala Pro Pro Ser
126 785          790          795          800
127 Ser Gly Gly Ser Arg Thr
128          805

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132 <212> TYPE: DNA
133 <213> ORGANISM: Artificial Sequence
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140 <211> LENGTH: 55
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142 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
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Input Set : N:\AMC\US10734661A.raw

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151 <213> ORGANISM: Homo sapiens
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153 <308> DATABASE ACCESSION NO: m58051
154 <309> DATABASE ENTRY DATE: 1994-11-08
155 <313> RELEVANT RESIDUES: (1)..(1147)
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159      agcagcgctg cgtggggcga ggggcagaag tccccggccc agagcccgcc cagcaggagc      180
160      agttggtctt cggcagcggg gatgctgtgg agctgagctg tcccccgccc gggggtggtc      240
161      ccatggggcc cactgtctgg gtcaaggatg gcacagggtg ggtgccctcg gagcgtgtcc      300
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168      atcagcagtg gagcctggtc atggaaagcg tggtgccctc ggaccgcggc aactacacct      720
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173      tgctcaagac ggcgggcgct aacaccaccg acaaggagct agaggttctc tccttgaca      1020
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181 <213> ORGANISM: Artificial sequence
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187      cgagcaaaat ttaagctaca acaaggcaag gcttgaccga caattgcatg aagaatctgc      180
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192      attgacgtca atgggtggac tatttacggg aaactgccca cttggcagta catcaagtgt      480
193      atcatatgcc aagtacgcc cctattgacg tcaatgacgg taaatggccc gcctggcatt      540
194      atgccagta catgacctta tgggactttc ctacttgga gtacatctac gtattagtca      600
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198	gtaggcgtgt	acgggtgggag	gtctatataa	gcagagctct	ctggctaact	agagaaccca	840
199	ctgcttactg	gcttatcgaa	attaatacga	ctcactatag	ggagacccaa	gctggctagc	900
200	gttttaaactt	aaagcttggt	ccgagctcgg	atccccgtcg	tgcattctatc	gaaggtcgtg	960
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202	ctgaactcct	ggggggaccg	tcagtcttcc	tcttcccccc	aaaacccaag	gacacctca	1080
203	tgatctcccg	gacccctgag	gtcacatgcg	tgggtggtgga	cgtgagccac	gaagacctg	1140
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205	gggaggagca	gtacaacagc	acgtaccggg	tggtcagcgt	cctcaccgtc	ctgcaccagg	1260
206	actggctgaa	tggcaaggag	tacaagtgc	aggtctccaa	caaagccctc	ccagccccc	1320
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209	tctatcccag	cgacatcgcc	gtggagtggg	agagcaatgg	gcagccggag	aacaactaca	1500
210	agaccacgcc	tcccgtgctg	gactccgacg	gctccttctt	cctctacagc	aagctcaccg	1560
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215	aaaatgagga	aattgcatcg	cattgtctga	gtaggtgtca	ttctattctg	gggggtgggg	1860
216	tggggcagga	cagcaagggg	gaggattggg	aagacaatag	caggcatgct	ggggatgcgg	1920
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239	cgtgctacga	gatttcgatt	ccaccgccc	cttctatgaa	aggttgggct	tcggaatcgt	3300
240	tttccgggac	gccggctgga	tgatcctcca	gcgcggggat	ctcatgctgg	agttcttcgc	3360
241	ccaccccaac	ttgtttattg	cagcttataa	tggttacaaa	taaagcaata	gcatacaaaa	3420
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:54; N Pos. 253,254,255
Seq#:56; N Pos. 256,257,258
Seq#:70; N Pos. 1,2,3
Seq#:74; N Pos. 1,2,3
Seq#:81; N Pos. 1,2,3
Seq#:83; N Pos. 1,2,3

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7
Seq#:7; Line(s) 326
Seq#:30; Line(s) 580
Seq#:31; Line(s) 589
Seq#:32; Line(s) 598
Seq#:33; Line(s) 607
Seq#:34; Line(s) 616
Seq#:35; Line(s) 625
Seq#:36; Line(s) 634
Seq#:37; Line(s) 643
Seq#:38; Line(s) 652
Seq#:39; Line(s) 661
Seq#:40; Line(s) 670
Seq#:41; Line(s) 679
Seq#:42; Line(s) 688
Seq#:43; Line(s) 697
Seq#:44; Line(s) 706
Seq#:45; Line(s) 715
Seq#:46; Line(s) 724
Seq#:47; Line(s) 733
Seq#:48; Line(s) 742
Seq#:49; Line(s) 751
Seq#:50; Line(s) 760
Seq#:51; Line(s) 769

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Input Set : N:\AMC\US10734661A.raw

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L:9 M:270 C: Current Application Number differs, Wrong Format

L:984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:240

L:1016 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:240

L:1212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0

L:1273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0

L:1380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:0

L:1413 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0